

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/567,938A
Source: IFWP
Date Processed by STIC: 11/15/2006

ENTERED



IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/567,938A

DATE: 11/15/2006

TIME: 11:14:36

Input Set : A:\211050011U2.txt

Output Set: N:\CRF4\11152006\J567938A.raw

4 <110> APPLICANT: Rowe, Peter S. N.
 6 <120> TITLE OF INVENTION: REGULATION OF TISSUE MINERALIZATION AND
 7 PHOSPHATE METABOLISM BY ASARM PEPTIDES
 10 <130> FILE REFERENCE: 21105.0011U2
 12 <140> CURRENT APPLICATION NUMBER: 10/567,938A
 C--> 13 <141> CURRENT FILING DATE: 2006-02-09
 15 <150> PRIOR APPLICATION NUMBER: PCT/US04/30530
 16 <151> PRIOR FILING DATE: 2004-09-20
 18 <150> PRIOR APPLICATION NUMBER: 60/504,044
 19 <151> PRIOR FILING DATE: 2003-09-19
 21 <160> NUMBER OF SEQ ID NOS: 24
 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 25 <210> SEQ ID NO: 1
 26 <211> LENGTH: 19
 27 <212> TYPE: PRT
 28 <213> ORGANISM: Homo sapien
 30 <400> SEQUENCE: 1
 31 Arg Asp Asp Ser Ser Glu Ser Ser Asp Ser Gly Ser Ser Ser Glu Ser
 32 1 5 10 15
 33 Asp Gly Asp
 36 <210> SEQ ID NO: 2
 37 <211> LENGTH: 18
 38 <212> TYPE: PRT
 39 <213> ORGANISM: Mus musculus
 41 <400> SEQUENCE: 2
 42 Arg Asp Ser Ser Glu Ser Ser Ser Gly Ser Ser Ser Glu Ser His
 43 1 5 10 15
 44 Gly Asp
 47 <210> SEQ ID NO: 3
 48 <211> LENGTH: 18
 49 <212> TYPE: PRT
 50 <213> ORGANISM: Rattus norvegicus
 52 <400> SEQUENCE: 3
 53 Arg Asp Ser Ser Glu Ser Ser Ser Gly Ser Ser Ser Glu Ser Ser
 54 1 5 10 15
 55 Gly Asp
 58 <210> SEQ ID NO: 4
 59 <211> LENGTH: 24
 60 <212> TYPE: PRT
 61 <213> ORGANISM: Homo sapien
 65 <400> SEQUENCE: 4
 66 Phe Ser Ser Arg Arg Asp Asp Ser Ser Glu Ser Ser Asp Ser Gly
 67 1 5 10 15

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Input Set : A:\211050011U2.txt
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68 Ser Ser Ser Glu Ser Asp Gly Asp
69 20
71 <210> SEQ ID NO: 5
72 <211> LENGTH: 25
73 <212> TYPE: PRT
74 <213> ORGANISM: Homo sapien
76 <400> SEQUENCE: 5
77 Cys Phe Ser Ser Arg Arg Arg Asp Asp Ser Ser Glu Ser Ser Asp Ser
78 1 5 10 15
79 Gly Ser Ser Ser Glu Ser Asp Gly Asp
80 20 25
82 <210> SEQ ID NO: 6
83 <211> LENGTH: 26
84 <212> TYPE: PRT
85 <213> ORGANISM: Homo sapien
87 <400> SEQUENCE: 6
88 Cys Gly Ser Gly Tyr Thr Asp Leu Gln Glu Arg Gly Asp Asn Asp Ile
89 1 5 10 15
90 Ser Pro Phe Ser Gly Asp Gly Gln Pro Phe
91 20 25
93 <210> SEQ ID NO: 7
94 <211> LENGTH: 5
95 <212> TYPE: PRT
96 <213> ORGANISM: Homo sapien
98 <400> SEQUENCE: 7
99 Ala Pro Thr Phe Gln
100 1 5
102 <210> SEQ ID NO: 8
103 <211> LENGTH: 5
104 <212> TYPE: PRT
105 <213> ORGANISM: Homo sapien
107 <400> SEQUENCE: 8
108 Asp Ser Glu Ser Ser
109 1 5
111 <210> SEQ ID NO: 9
112 <211> LENGTH: 5
113 <212> TYPE: PRT
114 <213> ORGANISM: Homo sapien
116 <400> SEQUENCE: 9
117 Ser Ser Ser Glu Ser
118 1 5
120 <210> SEQ ID NO: 10
121 <211> LENGTH: 15
122 <212> TYPE: PRT
123 <213> ORGANISM: Homo sapien
127 <400> SEQUENCE: 10
128 Ala Pro Thr Phe Gln Pro Gln Thr Glu Lys Thr Lys Gln Ser Cys
129 1 5 10 15
131 <210> SEQ ID NO: 11

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132 <211> LENGTH: 19
 133 <212> TYPE: PRT
 134 <213> ORGANISM: Homo sapien
 136 <400> SEQUENCE: 11
 137 Thr Asp Leu Gln Glu Arg Gly Asp Asn Asp Ile Ser Pro Phe Ser Gly
 138 1 5 10 15
 139 Asp Gly Gln
 142 <210> SEQ ID NO: 12
 143 <211> LENGTH: 19
 144 <212> TYPE: PRT
 145 <213> ORGANISM: Homo sapien
 147 <400> SEQUENCE: 12
 148 Gly Arg Gln Pro His Ser Asn Arg Arg Phe Ser Ser Arg Arg Arg Asp
 149 1 5 10 15
 150 Asp Ser Ser
 153 <210> SEQ ID NO: 13
 154 <211> LENGTH: 18
 155 <212> TYPE: PRT
 156 <213> ORGANISM: Homo sapien
 158 <400> SEQUENCE: 13
 159 Asp Asp Ser Ser Glu Ser Ser Asp Ser Gly Ser Ser Ser Glu Ser Asp
 160 1 5 10 15
 161 Gly Asp
 164 <210> SEQ ID NO: 14
 165 <211> LENGTH: 19
 166 <212> TYPE: PRT
 167 <213> ORGANISM: Homo sapien
 169 <220> FEATURE:
 170 <221> NAME/KEY: VARIANT
 171 <222> LOCATION: 12, 14, 16
 172 <223> OTHER INFORMATION: Xaa = a phosphorylated serine
 174 <400> SEQUENCE: 14
 W--> 175 Arg Asp Asp Ser Ser Glu Ser Ser Asp Ser Gly Xaa Ser Xaa Glu Xaa
 176 1 5 10 15
 177 Asp Gly Asp
 180 <210> SEQ ID NO: 15
 181 <211> LENGTH: 25
 182 <212> TYPE: PRT
 183 <213> ORGANISM: Homo sapien
 187 <400> SEQUENCE: 15
 188 Gly Ser Gly Tyr Thr Asp Leu Gln Glu Arg Gly Asp Asn Asp Ile Ser
 189 1 5 10 15
 190 Pro Phe Ser Gly Asp Gly Gln Pro Phe
 191 20 25
 193 <210> SEQ ID NO: 16
 194 <211> LENGTH: 19
 195 <212> TYPE: PRT
 196 <213> ORGANISM: Macaca fascicularis
 198 <400> SEQUENCE: 16

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199 Arg Glu Asp Ser Ser Glu Ser Ser Asp Ser Gly Ser Ser Ser Glu Ser
200 1 5 10 15
201 Asp Gly Asp
204 <210> SEQ ID NO: 17
205 <211> LENGTH: 525
206 <212> TYPE: PRT
207 <213> ORGANISM: Homo sapien
209 <400> SEQUENCE: 17
210 Met Arg Val Phe Cys Val Gly Leu Leu Phe Ser Val Thr Trp Ala
211 1 5 10 15
212 Ala Pro Thr Phe Gln Pro Gln Thr Glu Lys Thr Lys Gln Ser Cys Val
213 20 25 30
214 Glu Glu Gln Arg Gln Glu Glu Lys Asn Lys Asp Asn Ile Gly Phe His
215 35 40 45
216 His Leu Gly Lys Arg Ile Asn Gln Glu Leu Ser Ser Lys Glu Asn Ile
217 50 55 60
218 Val Gln Glu Arg Lys Lys Asp Leu Ser Leu Ser Glu Ala Ser Glu Asn
219 65 70 75 80
220 Lys Gly Ser Ser Lys Ser Gln Asn Tyr Phe Thr Asn Arg Gln Arg Leu
221 85 90 95
222 Asn Lys Glu Tyr Ser Ile Ser Asn Lys Glu Asn Thr His Asn Gly Leu
223 100 105 110
224 Arg Met Ser Ile Tyr Pro Lys Ser Thr Gly Asn Lys Gly Phe Glu Asp
225 115 120 125
226 Gly Asp Asp Ala Ile Ser Lys Leu His Asp Gln Glu Glu Tyr Gly Ala
227 130 135 140
228 Ala Leu Ile Arg Asn Asn Met Gln His Ile Met Gly Pro Val Thr Ala
229 145 150 155 160
230 Ile Lys Leu Leu Gly Glu Glu Asn Lys Glu Asn Thr Pro Arg Asn Val
231 165 170 175
232 Leu Asn Ile Ile Pro Ala Ser Met Asn Tyr Ala Lys Ala His Ser Lys
233 180 185 190
234 Asp Lys Lys Lys Pro Gln Arg Asp Ser Gln Ala Gln Lys Ser Pro Val
235 195 200 205
236 Lys Ser Lys Ser Thr His Arg Ile Gln His Asn Ile Asp Tyr Leu Lys
237 210 215 220
238 His Leu Ser Lys Val Lys Lys Ile Pro Ser Asp Phe Glu Gly Ser Gly
239 225 230 235 240
240 Tyr Thr Asp Leu Gln Glu Arg Gly Asp Asn Asp Ile Ser Pro Phe Ser
241 245 250 255
242 Gly Asp Gly Gln Pro Phe Lys Asp Ile Pro Gly Lys Gly Glu Ala Thr
243 260 265 270
244 Gly Pro Asp Leu Glu Gly Lys Asp Ile Gln Thr Gly Phe Ala Gly Pro
245 275 280 285
246 Ser Glu Ala Glu Ser Thr His Leu Asp Thr Lys Lys Pro Gly Tyr Asn
247 290 295 300
248 Glu Ile Pro Glu Arg Glu Glu Asn Gly Asn Thr Ile Gly Thr Arg
249 305 310 315 320
250 Asp Glu Thr Ala Lys Glu Ala Asp Ala Val Asp Val Ser Leu Val Glu

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251 325 330 335
 252 Gly Ser Asn Asp Ile Met Gly Ser Thr Asn Phe Lys Glu Leu Pro Gly
 253 340 345 350
 254 Arg Glu Gly Asn Arg Val Asp Ala Gly Ser Gln Asn Ala His Gln Gly
 255 355 360 365
 256 Lys Val Glu Phe His Tyr Pro Pro Ala Pro Ser Lys Glu Lys Arg Lys
 257 370 375 380
 258 Glu Gly Ser Ser Asp Ala Ala Glu Ser Thr Asn Tyr Asn Glu Ile Pro
 259 385 390 395 400
 260 Lys Asn Gly Lys Gly Ser Thr Arg Lys Gly Val Asp His Ser Asn Arg
 261 405 410 415
 262 Asn Gln Ala Thr Leu Asn Glu Lys Gln Arg Phe Pro Ser Lys Gly Lys
 263 420 425 430
 264 Ser Gln Gly Leu Pro Ile Pro Ser Arg Gly Leu Asp Asn Glu Ile Lys
 265 435 440 445
 266 Asn Glu Met Asp Ser Phe Asn Gly Pro Ser His Glu Asn Ile Ile Thr
 267 450 455 460
 268 His Gly Arg Lys Tyr His Tyr Val Pro His Arg Gln Asn Asn Ser Thr
 269 465 470 475 480
 270 Arg Asn Lys Gly Met Pro Gln Gly Lys Gly Ser Trp Gly Arg Gln Pro
 271 485 490 495
 272 His Ser Asn Arg Arg Phe Ser Ser Arg Arg Arg Asp Asp Ser Ser Glu
 273 500 505 510
 274 Ser Ser Asp Ser Gly Ser Ser Glu Ser Asp Gly Asp
 275 515 520 525
 277 <210> SEQ ID NO: 18
 278 <211> LENGTH: 433
 279 <212> TYPE: PRT
 280 <213> ORGANISM: Mus musculus
 282 <400> SEQUENCE: 18
 283 Met Gln Ala Val Ser Val Gly Leu Leu Leu Phe Ser Met Thr Trp Ala
 284 1 5 10 15
 285 Ala Pro Met Pro Asn Glu Asp Arg Ser Ser Cys Gly Asn Gln Asp Ser
 286 20 25 30
 287 Ile His Lys Asp Leu Ala Ala Ser Val Tyr Pro Asp Pro Thr Val Asp
 288 35 40 45
 289 Glu Gly Thr Glu Asp Gly Gln Gly Ala Leu Leu His Pro Pro Gly Gln
 290 50 55 60
 291 Asp Arg Tyr Gly Ala Ala Leu Leu Arg Asn Ile Thr Gln Pro Val Lys
 292 65 70 75 80
 293 Ser Leu Val Thr Gly Ala Glu Leu Arg Arg Glu Gly Asn Gln Glu Lys
 294 85 90 95
 295 Arg Pro Gln Ser Val Leu Ser Val Ile Pro Ala Asp Val Asn Asp Ala
 296 100 105 110
 297 Lys Val Ser Leu Lys Asp Ile Lys Asn Gln Glu Ser Tyr Leu Leu Thr
 298 115 120 125
 299 Gln Ser Ser Pro Val Lys Ser Lys His Thr Lys His Thr Arg Gln Thr
 300 130 135 140
 301 Arg Arg Ser Thr His Tyr Leu Thr His Leu Pro Gln Ile Lys Lys Thr

RAW SEQUENCE LISTING ERROR SUMMARY **DATE:** 11/15/2006
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:14; Xaa Pos. 12,14,16
Seq#:21; Xaa Pos. 1,2,3,4,6,11,13,14,15,16,17,19,20,21,22,29,30,80,92,93,94
Seq#:21; Xaa Pos. 95,96,97,98,99,106,107,110,111,112,114,115,116,117
Seq#:22; Xaa Pos. 1,6,8,9,10,11,12
Seq#:23; Xaa Pos. 7,8,11
Seq#:24; Xaa Pos. 46

VERIFICATION SUMMARY

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:562 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
M:341 Repeated in SeqNo=21
L:617 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:636 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:656 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:32